

AMENDMENTS**Amendments to the Claims:**

This listing of claims will replace all prior versions and listing of claims in the application:

Listing of Claims:

1. **(Previously presented)** A transgenic non-human animal, wherein said transgenic non-human animal comprises somatic or germline cells having incorporated into the genome of said cells a transgene that expresses an epitope-tagged TATA-box binding protein (TBP).
2. **(Previously presented)** The transgenic non-human animal of claim 1, wherein said epitope-tagged TBP comprises two epitope-tags.
3. **(Previously presented)** The transgenic non-human animal of claim 2, wherein said epitope-tagged TBP comprises human TBP (hTBP).
4. **(Previously presented)** The transgenic non-human animal of claim 3, wherein said two epitope-tags comprise an HA-epitope and a His-epitope.
5. **(Previously presented)** The transgenic non-human animal of claim 4, wherein said epitope-tagged TBP comprises the amino acid sequence of SEQ ID NO. 16.
6. **(Previously presented)** The transgenic non-human animal of claim 1, wherein the transgenic animal is a mouse.
7. **(Currently Amended)** A method of making a transgenic non-human animal, comprising the steps of:
 - (i) providing a transgene that expresses an epitope-tagged TATA-box binding protein;
 - (ii) introducing said transgene into ~~germline cells~~ an embryonic cell of said non-human animal, wherein said transgene is stably integrated into the genome of said cells cell;
 - (iii) transferring said transfected ~~germline cells~~ embryonic cell to a surrogate mother, and permitting said ~~germline~~-cell to develop into a non-human transgenic animal; and

- (iv) identifying a non-human transgenic animal that produces said epitope-tagged TATA-box binding protein.
8. **(Currently Amended)** The method of claim 7, wherein said transgene is introduced into said cell[[s]] by microinjecting said transgene into a zygote of said non-human animal.
9. **(Currently Amended)** The method of claim 7, wherein said transgene is introduced into said cell[[s]] by transfecting a blastomere of said non-human animal with a vector comprising said transgene.
10. **(Currently Amended)** The method of claim 7, wherein said transgene is introduced into said cell[[s]] by introducing said transgene into ~~an embryonic stem cell~~ a stem cell that is combined with a blastocyst of said non-human animal.
11. **(Previously presented)** The transgenic non-human animal of claim 1, wherein said transgene comprises a first DNA sequence that encodes for TBP and a second DNA sequence that encodes for one or more epitope-tags.
12. **(Previously presented)** The transgenic non-human animal of claim 11, wherein said first DNA sequence is a cDNA sequence.
13. **(Previously presented)** The transgenic non-human animal of claim 12, wherein said cDNA sequence is the cDNA of human TBP (hTBP).
14. **(Previously presented)** The transgenic non-human animal of claim 11, wherein said second DNA sequence encodes for two epitope-tags.
15. **(Previously presented)** The transgenic non-human animal of claim 14, wherein said two epitope-tags comprise an HA-epitope and a His-epitope.
16. **(Previously presented)** The transgenic non-human animal of claim 15, wherein said HA-epitope comprises an amino acid sequence selected from the group consisting of the amino acid sequences of SEQ ID NO. 1, SEQ ID NO. 2, SEQ ID NO. 3, and SEQ ID NO. 4.
17. **(Previously presented)** The transgenic non-human animal of claim 1, wherein said transgene comprises the nucleotide sequence of SEQ ID NO. 13.

18. **(Previously presented)** The transgenic non-human animal of claim 1, wherein said transgene further comprises an inducible or constitutive promoter.
19. **(Previously presented)** The transgenic non-human animal of claim 18, wherein said constitutive promoter is the promoter of an elongation factor-1 alpha gene (EF) gene.
20. **(Previously presented)** The transgenic non-human animal of claim 18, wherein said inducible promoter is the promoter of an metallothionine promoter (MT) gene.
21. **(Previously presented)** The transgenic non-human animal of claim 19, wherein said transgene comprises the nucleotide sequence of SEQ ID NO. 14.
22. **(Previously presented)** The transgenic non-human animal of claim 20, wherein said transgene comprises the nucleotide sequence of SEQ ID NO. 15.
23. **(Previously presented)** A method of expressing an epitope-tagged TBP in a non-human transgenic animal, comprising introducing a transgene encoding said epitope-tagged TBP into cells selected from the group consisting of:
 - (a) germline cells of said non-human animal;
 - (b) somatic cells of said non-human animal; and
 - (c) both germline cells and somatic cells of said non-human animal wherein said epitope tagged TBP is expressed in said non-human transgenic animal.
24. **(Previously presented)** The method of claim 23, wherein said transgene further comprises a constitutive promoter.
25. **(Previously presented)** The method of claim 23, wherein said transgene further comprises an inducible promoter and wherein said epitope-tagged TBP is expressed upon induction of said promoter.
26. **(Previously presented)** A method for isolating a higher order transcription complex, comprising introducing a transgene encoding an epitope-tagged TBP into a non-human animal, expressing said epitope-tagged TBP in said animal, wherein said epitope-tagged TBP binds to one or more TBP-associated factors (TAFs) to create said higher order transcription complex, and isolating said higher order transcription complex from said animal.

27. **(Previously presented)** The method of claim 26, wherein said higher order transcription complex further comprises a TAF-interacting factor bound to said one or more TAFs.
28. **(Previously presented)** The method of claim 26, further comprising isolating said higher order transcription complex from more than one type of tissue or more than one cell type of said transgenic non-human animal.
29. **(Previously presented)** The method of claim 26, further comprising isolating said higher order transcription complex from said transgenic non-human animal during two or more different developmental stages of said animal.
30. **(Previously presented)** A method for isolating a TAF or a TAF-interacting factor, comprising introducing a transgene encoding an epitope-tagged TBP into a non-human animal, expressing said epitope-tagged TBP in said animal wherein said epitope-tagged TBP binds to said TAF or TAF-interacting factor to form a complex, isolating said complex, and separating said TAF or TAF-interacting factor from said epitope-tagged TBP.
31. **(Previously presented)** The method of claim 30, further comprising determining the amino acid sequence of said TAF or TAF-interacting factor.
32. **(Previously presented)** The method of claim 26, wherein said higher order transcription complex is isolated by contacting said complex with an affinity purification column comprising, as a solid phase, antibodies that bind to at least one of the epitopes of said epitope-tagged TBP component of said higher order transcription complex.
33. **(Previously presented)** The method of claim 32, wherein at least one epitope of said epitope-tagged TBP is a His epitope and wherein said affinity purification column is a Ni^{2+} .
34. **(Previously presented)** The method of claim 32, wherein at least one epitope of said epitope-tagged TBP is an HA epitope and wherein said affinity purification column comprises, as a solid phase, anti-HA antibodies.
35. **(Previously presented)** The method of claim 32, wherein at least one epitope of said epitope-tagged TBP comprises the amino acid sequence of SEQ ID NO. 17 and wherein said affinity purifying column comprises, as a solid phase, antibodies that bind to said epitope.

36. **(Canceled)** An antibody that binds to an epitope-tagged TBP, wherein said epitope comprises the amino acid sequence of SEQ ID NO. 17.
37. **(Previously presented)** The transgenic non-human animal of claim 4, wherein said HA epitope comprises an amino acid sequence selected from the group consisting of SEQ ID NO. 1, SEQ ID NO. 2, SEQ ID NO. 3 and SEQ ID NO. 4.